

Substitute SEQUENCE LISTING

<110> Kwon, Byoung

<120> NEW RECEPTOR AND RELATED PRODUCTS AND METHODS

<130> 740.013US2

<140> 08/955,572

<141> 1997-10-22

<150> 08/461,652

<151> 1995-06-05

<150> 08/122,796

<151> 1993-09-03

<160> 12

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 838

<212> DNA

<213> Homo sapiens

<400> 1

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ttgtagtaac	tgcccagctg	gtacattctg	tgataataac	aggaatcaga	tttgagctcc	180
ctgtcctcca	aatagtttct	ccagcgcagg	tgacaaaagg	acctgtgaca	tatgcaggca	240
gtgtaaaggt	gttttcagga	ccaggaagga	gtgttcctcc	accagcaatg	cagagtgtga	300
ctgcactcca	gggttttact	gcctgggggc	aggatgcagc	atgtgtgaac	aggattgtaa	360
acaaggtcaa	gaactgacaa	aaaaagggtg	taaagactgt	tgctttggga	catttaacga	420
tcagaaacgt	ggcatctgtc	gacctgggac	aaactgttct	ttggatggaa	agtctgtgct	480
tgtgaatggg	acgaaggaga	gggacgtggt	ctgtggacca	tctccagctg	acctctctcc	540
gggagcatcc	tctgtgaccc	cgccctgccc	tgcgagagag	ccaggacact	ctccgcagat	600
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gctccgtttc	tctgttggtt	aacggggcag	aaagaaaactc	ctgtatatat	tcaaacaacc	720
atztatgaga	ccagtacaaa	ctactcaaga	ggaagatggc	tgtagctgcc	gatttccaga	780
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<210> 2

<211> 255

<212> PRT

<213> Homo sapiens

<400> 2

Met	Gly	Asn	Ser	Cys	Tyr	Asn	Ile	Val	Ala	Thr	Leu	Leu	Leu	Val	Leu
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Asn	Phe	Glu	Arg	Thr	Arg	Ser	Leu	Gln	Asp	Pro	Cys	Ser	Asn	Cys	Pro
			20					25					30		
Ala	Gly	Thr	Phe	Cys	Asp	Asn	Asn	Arg	Asn	Gln	Ile	Cys	Ser	Pro	Cys
			35				40					45			
Pro	Pro	Asn	Ser	Phe	Ser	Ser	Ala	Gly	Gly	Gln	Arg	Thr	Cys	Asp	Ile
		50				55				60					
Cys	Arg	Gln	Cys	Lys	Gly	Val	Phe	Arg	Thr	Arg	Lys	Glu	Cys	Ser	Ser
65				70				75						80	
Thr	Ser	Asn	Ala	Glu	Cys	Asp	Cys	Thr	Pro	Gly	Phe	His	Cys	Leu	Gly

<210> 8
 <211> 30
 <212> DNA
 <213> Homo sapiens

<400> 8
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30

<210> 9
 <211> 2350
 <212> DNA
 <213> Mus musculus

<220>
 <221> unsure
 <222> (1253)...(1255)
 <223> (a or g or c or t/u)

<400> 9

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ggacccccacc	atcctgtgga	acagcacaa	caacccccacc	accctgttct	tacacatcat	1020
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acacacacac	acacacacac	acacacacgt	ttatactacg	tactgttata	ggtattctac	1560
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atctcacaag	tttcgtccgg	gtcggcgga	cctatggcgt	cgatccttat	taccttatcc	1980
tggcgccaag	ataaaacaac	caaaagcctt	gactccggta	ctaattctcc	ctgccggccc	2040
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ctttcgtaaa	cggttcttac	aaaagtaatt	agttcttgct	ttcagcctcc	aagcttctgc	2160
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<210> 10

<211> 256
 <212> PRT
 <213> Mus musculus

<400> 10
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 1 5 10 15
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 Pro Gly Thr Phe Cys Arg Lys Tyr Asn Pro Val Cys Lys Ser Cys Pro
 35 40 45
 Pro Ser Thr Phe Ser Ser Ile Gly Gly Gln Pro Asn Cys Asn Ile Cys
 50 55 60
 Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser Thr
 65 70 75 80
 His Asn Ala Glu Cys Glu Cys Ile Glu Gly Phe His Cys Leu Gly Pro
 85 90 95
 Gln Cys Thr Arg Cys Glu Lys Asp Cys Arg Pro Gly Gln Glu Leu Thr
 100 105 110
 Lys Gln Gly Cys Lys Thr Cys Ser Leu Gly Thr Phe Asn Asp Gln Asn
 115 120 125
 Gly Thr Gly Val Cys Arg Pro Trp Thr Asn Cys Ser Leu Asp Gly Arg
 130 135 140
 Ser Val Leu Lys Thr Gly Thr Thr Glu Lys Asp Val Val Cys Gly Pro
 145 150 155 160
 Pro Val Val Ser Phe Ser Pro Ser Thr Thr Ile Ser Val Thr Pro Glu
 165 170 175
 Gly Gly Pro Gly Gly His Ser Leu Gln Val Leu Thr Leu Phe Leu Ala
 180 185 190
 Leu Thr Ser Ala Leu Leu Leu Ala Leu Ile Phe Ile Thr Leu Leu Phe
 195 200 205
 Ser Val Leu Lys Trp Ile Arg Lys Lys Phe Pro His Ile Phe Lys Gln
 210 215 220
 Pro Phe Lys Lys Thr Thr Gly Ala Ala Gln Glu Glu Asp Ala Cys Ser
 225 230 235 240
 Cys Arg Cys Pro Gln Glu Glu Glu Gly Gly Gly Gly Tyr Glu Leu
 245 250 255

<210> 11
 <211> 24
 <212> PRT
 <213> Homo sapiens

<220>
 <221> ZN_FING
 <222> 2...3, 5...13, 15...17, 19...21, 23
 <223> Putative zinc finger structure

<400> 11
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 1 5 10 15
 Xaa His Xaa Xaa Xaa Cys Xaa Cys
 20

<210> 12
 <211> 12
 <212> PRT
 <213> Homo sapiens

<400> 12

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